

FIG.1

Fas/AP01
Ankyrin1
p55 TNF-R
NGF-R
MORT1

VKGFVAKNGVAKIDETIKNDNVQDTAEQKVQLLRNWHQHGKKHA.YPTLTKIDKKKANDCTDA 307
WAEIARELOFSVEDINRIRVENPNSITAEQSVALLNIMVIREGONANMENHVTALQSIDRGEIV 1479
WKEFVRLGLSDHEIDFLELQNGRQLAEQAQYSMLATMRRTRPRREATLEILGRVLRDHLGLCL 434
WRHLAGELGYPPEHDSFTHE...ACPVRALLASMA...TQDSATLDALLALRRIORALIV 414
MRRLARLRVSDTKIDSIEDRYPRNTAEVRVRESLRIMKN.TEKENATVAHLVGAIRSCCMN LV 210

FIG. 2

DBD CONSTRUCT	AD CONSTRUCT	SCORE	<i>lpr</i> ^{cg} -like mutation effect		
			IN DBD CONSTRUCT	IN AD CONSTRUCT	IN BOTH
p55	p55				
Fas	Fas				
TRADD	TRADD		ND	ND	ND
MORT 1	MORT 1				
RIP	RIP				
TRADD	p55		ND		ND
p55	TRADD			ND	ND
TRADD	Fas		ND		ND
Fas	TRADD			ND	ND
MORT 1	p55				
p55	MORT 1				
MORT 1	Fas				
Fas	MORT 1				
RIP	p55				
p55	RIP				
RIP	Fas				
Fas	RIP				
TRADD	MORT 1		ND		ND
MORT 1	TRADD			ND	ND
TRADD	RIP		ND		ND
RIP	TRADD			ND	ND
MORT 1	RIP				
RIP	MORT 1				

<1h

1-2h

2-24h

NEGATIVE

FIG. 3

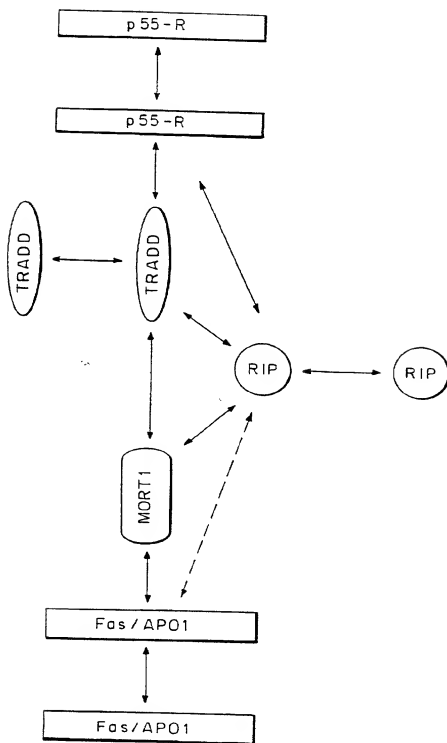


FIG. 4

